

Amendments to the Claims:

This listing of claims will replace all prior versions, and listings, of claims in the application.

Listing of Claims:

1-4. (canceled)

1/5. (currently amended) A method of diagnosing hepatocellular carcinoma in a human patient, comprising:

(a) detecting the level of expression in a tissue sample of two or more genes from Tables 3A, 3B, 5, 7A, 7B, 8A and 8B [3-9]; wherein differential expression of the genes in Tables 3A, 3B, 5, 7A, 7B, 8A and 8B [3-9] is indicative of hepatocellular carcinoma.

2/6. (currently amended) A method of detecting the progression of hepatocellular carcinoma in a human patient, comprising:

(a) detecting the level of expression in a tissue sample of two or more genes from Tables 3A, 3B, 5, 7A, 7B, 8A and 8B [3-9]; wherein differential expression of the genes in Tables 3A, 3B, 5, 7A, 7B, 8A and 8B [3-9] is indicative of hepatocellular carcinoma progression.

3/7. (currently amended) A method of monitoring the treatment of a human patient with hepatocellular carcinoma, comprising:

- (a) administering a pharmaceutical composition to the patient;
- (b) preparing a gene expression profile from a cell or tissue sample from the patient; and
- (c) comparing the patient gene expression profile to a gene expression from a cell population comprising normal liver cells or to a gene expression profile from a cell population comprising hepatocellular carcinoma cells or to both.

8. (canceled)

9. (currently amended) A method of diagnosing a metastatic liver [tumor] cancer in a human patient, comprising:

(a) detecting the level of expression in a tissue sample of two or more genes from Tables 4A, 4B, 5, 6A, 6B, 9A and 9B [3-9]; wherein differential expression of the genes in Tables 4A, 4B, 5, 6A, 6B, 9A and 9B [3-9] is indicative of metastatic liver cancer.

10. (currently amended) A method of detecting the progression of a metastatic liver cancer in a human patient, comprising:

(a) detecting the level of expression in a tissue sample of two or more genes from Tables 4A, 4B, 5, 6A, 6B, 9A and 9B [3-9]; wherein differential expression of the genes in Tables 4A, 4B, 5, 6A, 6B, 9A and 9B [3-9] is indicative of a metastatic liver cancer progression.

11. (currently amended) A method of monitoring the treatment of a human patient with a metastatic liver cancer, comprising:

- (a) administering a pharmaceutical composition to the patient;
- (b) preparing a gene expression profile from a cell or tissue sample from the patient; and
- (c) comparing the patient gene expression profile to a gene expression from a cell population comprising normal liver cells or to a gene expression profile from a cell population comprising metastatic liver tumor cells or to both.

12. (canceled)

13. (currently amended) A method of differentiating metastatic liver cancer from hepatocellular carcinoma in a human patient, comprising:

(a) detecting the level of expression in a tissue sample of two or more genes from Tables 3-9; wherein differential expression of the genes in Tables 3-9 is indicative of metastatic liver cancer rather than hepatocellular carcinoma.

14 - 46. (canceled)

8 ~~47~~. (currently amended) A method of claim [1] ^{1 A} ~~5~~ or ~~9~~, wherein the level of expression of 5 or more genes from [Tables 3-9] one or more tables selected from the group consisting of Tables 3A, 3B, 4A, 4B, 5, 6A, 6B, 7A, 7B, 8A, 8B, 9A and 9B is detected.

9 ~~48~~. (currently amended) A method of claim [1] ^{1 A} ~~5~~ or ~~9~~, wherein the level of expression of 10 or more genes from [Tables 3-9] one or more tables selected from the group consisting of Tables 3A, 3B, 4A, 4B, 5, 6A, 6B, 7A, 7B, 8A, 8B, 9A and 9B is detected.

10 ~~49~~. (currently amended) A method of claim [1] ^{1 A} ~~5~~ or ~~9~~, wherein the level of expression of 100 or more genes from [Tables 3-9] one or more tables selected from the group consisting of Tables 3A, 3B, 4A, 4B, 5, 6A, 6B, 7A, 7B, 8A, 8B, 9A and 9B is detected.

11 ~~50~~. (currently amended) A method of claim [1] ^{1 A} ~~5~~ or ~~9~~, wherein the level of expression is compared to the gene information in [Tables 3-9] one or more tables selected from the group consisting of Tables 3A, 3B, 4A, 4B, 5, 6A, 6B, 7A, 7B, 8A, 8B, 9A and 9B.

12 ~~51~~. (currently amended) A method of diagnosing [liver cancer] hepatocellular carcinoma or metastatic liver cancer in a human patient comprising:

- (a) preparing a gene expression profile from a tissue sample; and
- (b) comparing the gene expression profile to a database comprising part of the data in Tables 3-9.

13 ~~52~~. (previously presented) A method of claim ¹² ~~51~~, wherein the cancer is hepatocellular carcinoma.

14 ~~53~~. (currently amended) A method of claim ¹² ~~51~~, wherein the cancer is metastatic [lung] liver cancer.

54. (canceled)

¹⁵~~55~~. (previously presented) A method of claim ¹²~~51~~, wherein the database comprises all of the data from Tables 3-9.

~~56~~. (canceled)

¹⁶~~57~~. (previously presented) A method of claim ¹²~~51~~, wherein the database comprises gene expression information for all of the genes from Tables 3-9.

¹⁷~~58~~. (new) A method of diagnosing hepatocellular carcinoma in a human patient, comprising:

(a) detecting the level of expression in a liver tissue sample of two or more mRNA species from Table 3A, 3B, 5, 7A, 7B, 8A or 8B; and

(b) comparing the detected level of expression to the level of expression of said two or more mRNA species in a hepatocellular carcinoma liver tissue sample, thereby diagnosing hepatocellular carcinoma in the patient.

¹⁸~~59~~. (new) A method of diagnosing metastatic liver cancer in a human patient, comprising:

(a) detecting the level of expression in a liver tissue sample of two or more mRNA species from Table 4A, 4B, 5, 6A, 6B, 9A or 9B; and

(b) comparing the detected level of expression to the level of expression of said two or more mRNA species in a metastatic liver cancer tissue sample, thereby diagnosing metastatic liver cancer in the patient.

¹⁹~~60~~. (new) A method of claim ¹⁷~~58~~, wherein the level of expression of said 2 or more mRNA species in a hepatocellular carcinoma liver sample is in Table 3A, 3B, 5, 7A, 7B, 8A or 8B.

²⁰~~61~~. (new) A method of claim ¹⁸~~59~~, wherein the level of expression of said 2 or more mRNA species in a metastatic liver cancer sample is in Table 4A, 4B, 5, 6A, 6B, 9A or 9B.

REMARKS